

**IN THE CLAIMS**

Please AMEND the claims as follows:

1. (Currently Amended) A soybean seed exhibiting an oil composition comprising 55 to 80% by weight oleic acid, ~~10 to 40% by weight linoleic acid, 6% or less by weight linolenic acid,~~ and 2 to 8% by weight saturated fatty acids, wherein said soybean seed further comprises a genome with a nucleic acid sequence that suppresses the expression of endogenous soybean *FAD2-1A*, comprising a fragment at least 95% identical to a soybean *FAD2-1A* intron that is at least 100 nucleotides in length; and said genome further comprises a nucleic acid sequence that increases the expression of beta-ketoacyl-ACP synthase IV.

2. (Canceled)

3. (Currently Amended) The soybean seed of claim ~~2~~ 1, wherein said seed exhibits an increased oleic acid content, a reduced saturated fatty acid content, and a reduced polyunsaturated fatty acid content relative to seed from a plant with a similar genetic background but lacking the recombinant nucleic acid molecule.

4. (Currently Amended) The soybean seed of claim ~~2~~ 1, wherein the oil composition further comprises 10 to 39% by weight linoleic acid~~[[,]]~~ and 4.5 ~~10%~~ 10% or less by weight linolenic acid, ~~and 3 to 6% by weight saturated fatty acids.~~

5. (Currently Amended) The soybean seed of claim ~~2~~ 1, wherein the oil composition further comprises 10 to 39% by weight linoleic acid~~[[,]]~~ ~~3-9~~ 3-9 ~~and 8.0%~~ and 8.0% or less by weight linolenic acid, ~~and 2 to 3.6% by weight saturated fatty acids.~~

6. (Currently Amended) The soybean seed of claim ~~2~~ 1, wherein the oil composition further comprises 11 to 30% by weight linoleic acid, ~~4.5~~ 6.0% ~~or less by weight linolenic acid,~~ and less than 6% by weight saturated fatty acids.

7 - 8. (Canceled)

9. (Currently Amended) A container of soybean seeds, wherein at least 25% of the seeds exhibit an oil composition comprising 55 to 80% by weight oleic acid, ~~10 to 40% by weight linoleic acid, 6% or less by weight linolenic acid,~~ and 2 to 8% by weight saturated fatty acids, wherein said soybean seeds further comprises a genome with a nucleic acid sequence that suppresses the expression of endogenous soybean *FAD2-1A*, comprising a fragment at least 95% identical to a soybean *FAD2-1A* intron that is at least 100 nucleotides in length; and said genome further comprises a nucleic acid sequence that increases the expression of beta-ketoacyl-ACP synthase IV.

10. (Currently Amended) A soybean seed exhibiting an oil composition comprising 65 to 80% by weight oleic acid, ~~10 to 30% by weight linoleic acid, 6% or less by weight linolenic acid,~~ and 2 to 8% by weight saturated fatty acids, wherein said soybean seed further comprises a genome with a nucleic acid sequence that suppresses the expression of endogenous soybean *FAD2-1A*, comprising a fragment at least 95% identical to a soybean *FAD2-1A* intron that is at least 100 nucleotides in length; and said genome further comprises a nucleic acid sequence that increases the expression of beta-ketoacyl-ACP synthase IV.

11. (Currently Amended) The soybean seed of claim 10, wherein the oil composition further comprises 10 to 29% by weight linoleic acid[[,]] ~~4-5~~ and 10% or less by weight linolenic acid, ~~and 3 to 6% by weight saturated fatty acids.~~

12. (Currently Amended) The soybean seed of claim 10, wherein the oil composition further comprises 10 to 29% by weight linoleic acid, ~~3-8~~ 8% or less by weight linolenic acid, and 2 to 3.6% by weight saturated fatty acids.

13-21. (Canceled)

22. (Currently Amended) A transformed soybean plant bearing seed, wherein said seed exhibits an oil composition which comprises 55 to 80% by weight oleic acid, ~~10 to 40% by weight linoleic acid, 6% or less by weight linolenic acid,~~ and 2 to 8% by weight saturated fatty

acids, wherein said seed further comprises a genome with a nucleic acid sequence that suppresses the expression of endogenous soybean *FAD2-1A*, comprising a fragment at least 95% identical to a soybean *FAD2-1A* intron that is at least 100 nucleotides in length; and said genome further comprises a nucleic acid sequence that increases the expression of beta-ketoacyl-ACP synthase IV.

23-24. (Canceled)

25. (Currently Amended) A plant part derived from the transformed plant of claim ~~23~~ 22.

26. (Currently Amended) Seed derived from the transformed plant of claim ~~23~~ 22.

27-72. (Canceled)